	Total	N-Clones (208F-FE-8)	T-Clones (FE8-208F)
Number of sequenced cDNA clones	1257	669	588
Number of individual sequences	823	416	407
Sequence analysis			
Known genes (nr/Genbank)	427	207	220
Expressed Sequence Tags (dbest)	303	161	142
No similarity in data bases (new)	93	48	45
Expression analysis: Reverse Northern Analysis/con- ventional Northern Blot			
Differentially expressed	393	225	168
Known genes	244	126	118
Expressed sequence tags	104	74	3 0
New sequences	45	25	20
Not differentially expressed	194	86	108
Not detectable in expression analysis	236	105	131

FIG. 1

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(;; 	Л		eri Eri		CAXI profein-tyrosine kinase	١.	3.7.2.356		£-
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# FIG. 2A

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FIG. 2B

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### Extracellular Profesins

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## FIG. 2C

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						Sery_ tRNA synthetase	900837 U	۹	 		
						Synexin (annexin VII)	E 113129	5.3	2.2		
							STA C	AF090791	2.3		
						TSG18: (tund pasceptibility protect)	276251 -	0.00	2.2		
						Tyrosine phosphatase-like protein 1A-2a; P1935	r 740652	552	74.	9 IT4,8	č.
											Ņ

Others

#### FIG. 2D

Expressed Sequence Tags (EST)

ESTS																		58000
Up-adjusted ESIS	8-213-04-50-28 8-3-1-04-60-28	ESTAB945139 ESTA mij7739	ESTATU31015	F.S.A.1.044101 F.S.A.1.7345.75	ES:076796	ESTHSACOUID/O	ES.WZ0810	LS: N65969										20 New Seguences
	F. A. A. 1499	FRIABLESUDS FRIABA	h.s.A.B. 1.44	5 5 5 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7	ES: AA 142426	FELTAL SURGES	F. AARTING	F.S. AAB. 3.5	ESTAM84 689	ESTAA850112	ESTAM350.23	ESTABA43333	15188314318	FELAA859425	ESTAA831266	1000 FEET 1000	New Seggerters	
	2 - FAM - 16 11 5 - 1 A A K + 1 4 6 4 6	ESTABATORE	LalAABaqûsî	1 1 AAAA 2 1 2 1 3 4 1 4 1 4 1 4 1 4 1 4 1 4 1 4 1 4 1	55 (AA893976	Es PAAguujug	2 × 1 AAA999584	ES!AA900577	ESTAM901340	£0TAA924035	r.STAA486886	P. STAIL SURGA	£51435777	13,1497,388			S. WOX.	
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COWC-ad	1. 1844. 36 1 3 2 × 1844. 7 18 1. 18	XCL TXXXXXX	ESTAA372927	17/25×44/51	cs:AA462855	25.7444.7442	±51AA51F974	ES19A617263	FSTAA517339	53 TAA5 72112	ESTAA5 (5650)	ENTARA SANTE	FSTAARSTS, 3	-STAAngello	ESTAA667812	1512A614131		Max 47
	3. M. 1.80	ESTANCE 33320	COCCOCALCE	25 An	33.AA.34455	38141£1894	1STAA. 03325	55550 THE 25	48.334.77.629	15TAA400432	44	75. AA. 47303	GORAC THATES	12 Me. 6 14	78.7A42×8366	SPACE SPACE		

## FIG. 2E

 $x^{(1)}$ 

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(JOSEPH COMPANY)	2337	11.00	FE3	Gerbaniv (Watt)	7000	10	(0) [1]
			-FC				<u>6</u> .
Grayacaky Graenyiqlutary, coa synthase	÷		÷	Bleomycin nyarolase		:	:
ABP-283 (actin binding protein/filamin)	÷	:+	+ + +	BRCAl-associated RING protein (Rand))	O	÷	
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APPS (Acentaringphenetical production)	+	< 5	<u>+</u>	FEW-1 (flap endonuclease-1)	O	+++++++++++++++++++++++++++++++++++++++	ŧ
73.5	-	63	++++	FKRP11 (T-Lell-Specific immunopaile)	15	;	
Controllers pusheun Cavell (a)	÷	<b>C</b> 5		Fire (FILSE-like likiblony protess)	್ಯು	;	c >
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Caffel (Agiory stammalating factor )	<u>;</u>	<:	:	TAPIC ('amina assettated polypeptide in	(,)	:	80
The total of the second of the	:	¢ >	÷	MAM domain protein	53	:	
FRS. transcription factor	<u>+</u>	+	÷	MAP-Kinase phosphatase (opg21) (c)	1.5	:	٠
TOLOGIA TOT COMPOSE AT LETE	;	6.5	<b>:</b>	$XXP = \{\emptyset \mid (Stronelysin = Z\} \mid (d)\}$	13	:	· ·
\$1.00 C	÷	+	<u>+</u>	MMP-3 (Stromelysin-1)	£2.	÷	<b>C</b> >
Solisatin-related protein, 19006			÷	Myb-binding protein (P160)	٠	:	,
STATE OF THE STATE	:	+	<u>+</u>	NF-1 transcription factor	<b></b> >	į	( >
11-0-10-10-10-10-10-10-10-10-10-10-10-10	;	c ,	-40	Not-regional enolgse (NNE)		:	÷
Best shook prohess 93	÷	es.		ORALIN (150 NO A PRYGAN PAGULATED)	٠	:	•
See seed firstegipting toyanders.	:	C	÷	per (samples) for the profess)	(3	;	:
interferon induced gene	-+	€.	.:	PKB 4 Mase	0	++++	+
li retropóson (ORF2)	:	< h	÷ ÷	RapiB GIP binding protein (A)	0	÷ ÷	,
270 200 200 200 200 200 200 200 200 200	÷	+	<b>‡</b>	Ras-GIPase-activating protein	<>	+++++++++++++++++++++++++++++++++++++++	,
	:	Ç.	-	Recal (rut spinocorobellar ataxia yene)	С	++++	+
Lysyl oxidase-related pratein (WM9-14)	+	C.		SA 1 (stromal antigen)	0	;	+
Mana dene	÷	$\Diamond$	+	Sortl (Sertilin)	0	++	; <del>+</del>
MMP-2 (Gelatinase A)	-	0	+++	TSG101 (tumor susceptibility protein)	÷	+++	+
mIEE3 (transcriptional activator)	+	+	<u>;</u>				
Nuclear autoantigen GS2NA	+	0	<u>+</u>				
Osteoglycip	+	0	÷				
PS protein	+++	+	÷				
בייבטבס-ק	÷ ;÷	<b>c</b> >	+				
Phosoduluthike protein (Phile)	+	0	+				
Serum indusible kinase (SMK)	+	()	<b>.</b>				
STATES CHAINCE SOLD FOLLOW	++	<>	+ + +				
Thrombospondin 1	+++++	O	+				
TIMP-2 (inhibitor of metalloproteinase 2)	÷	+	<b>:</b>				
785V-7/0115 pr. r (b)	+++++++++++++++++++++++++++++++++++++++	+	<b>†</b>				

### FIG. 3

	Ex	pressio	n Stre	ngth
Sequence Identity (Genbank/EMBL)	208F	FE-8 H-Ras	208F K-Ras	
ABC transporter MOAT-B	5,	++++	(	•
BDSD-1 (breast cancer suppressor candidate 1)	•	++++	(	-
Cyplooxygenase 1	+-	++++	+	++++
ElB 19R:Bol-2-binding protein (Nip3)	Ĵ	++	++++	++
EST AA743557	+++-	+	(	++
EST AA792426	4	++++	+	÷
EST AA904000	-	++++	+	+ +
ETF TEA aimain containing transcription factor	++++	C	+ +	+ +
Hamesyl pyrophosphate synthetase	-40	+++	C	÷-
FEM-1 (: lap endonuclease-1)	j	++++	+	3
FLIF (FLACE-like inhibitory protein)	7.0	+	+ +	-++-
CAP1 protein tyrosine kinase 1	_	++++	+	-+
MAGE-B dene cluster	÷.	++++	O	Ĉ
MAT-kinase phosphatase (opg21)		± +	+++	-++-
MARCKS	++++	('	+	+ + +
NMF-10 (Stromelysin 2)	j.	++	+ +	++++
$\operatorname{MoB}(-1)$ (F)	.:	+++;	+	-+
mTFE: (M-linked transcriptional activator)	++++	(	+	-1
MyN-binaing grotein (P16])		++++	+ -	++
nomel transcript N317	-+++	(	+ -	+++-
F-gadherin (a)	++++	(	C.	++
Phosphatidylinositol 3-kinase p170	+ + +	(;	+	+ +
Ran-GTPase-activating protein	Ç.	++++	C	0
SBM1 phospharase	Ü,	++++	+	+
Secum induciple kinase (SNK) (h)	++++	(ı	+++	+++
Typosine phosphatase IA-2a $(i)$	0	++++	С	++

FIG. 4

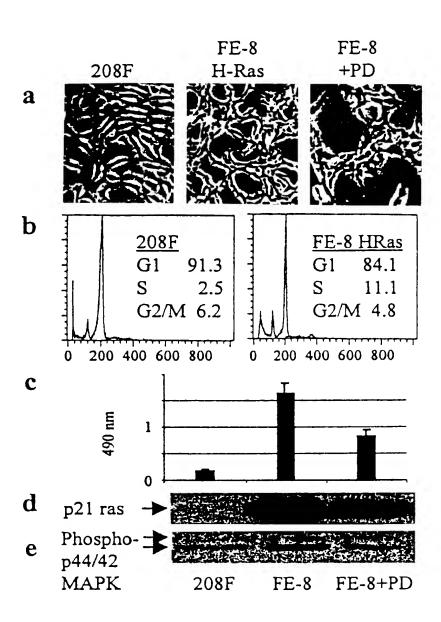


FIG. 5

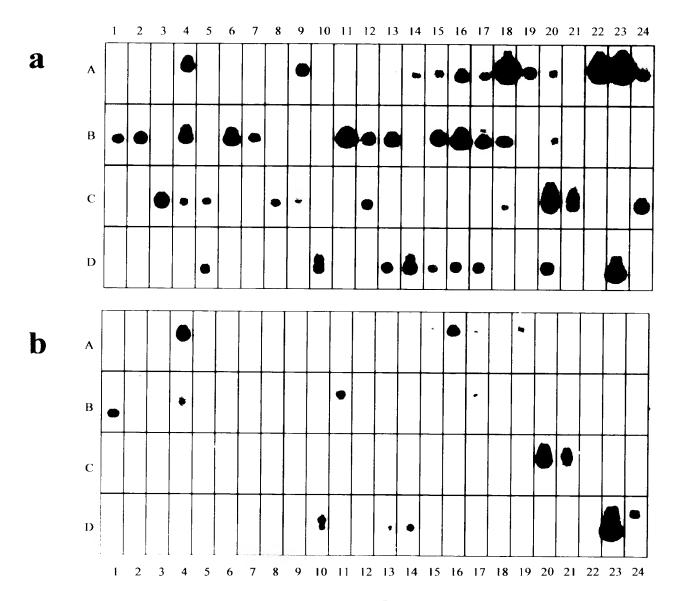
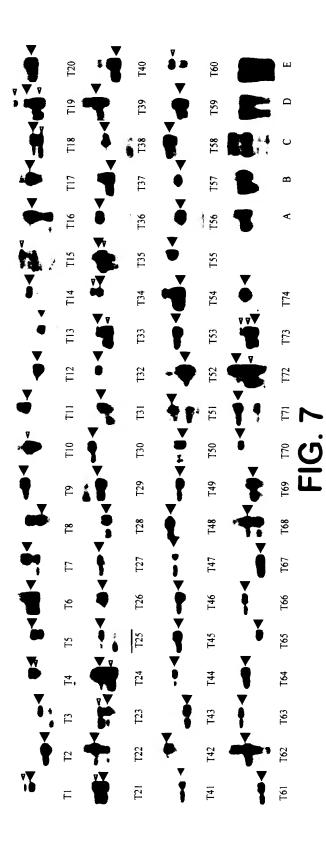


FIG. 6

NI6 NI7 NI8 NI9 N20 98 N N32 N33 N34 N35 N36 N37 N38 N35 N29 11 N58 N57 ? ?! N56 N9 NIO NII NI2 NI3 NI4 NI5 N53 N54 N55 1 1 N52 N51 N2S N26 N27 N28 N29 N30 N31 1 N44 N45 N46 N47 N48 N49 N50 02N 69N 7 N61 N62 N63 N64 N65 N66 N67 N68 % 1 \* \* Z 7 N2 N3 N4 N5 N6 \*\* V24 N41 N42 N43 Z2N Š ī



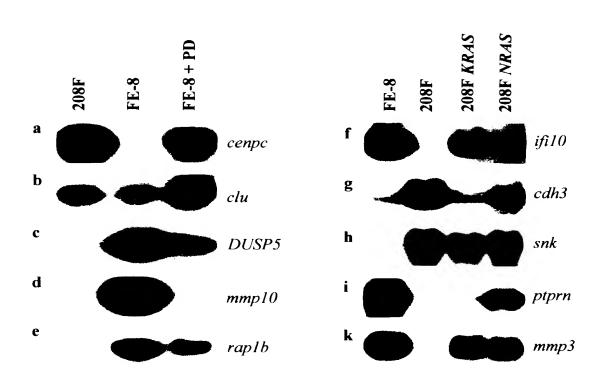


FIG. 8

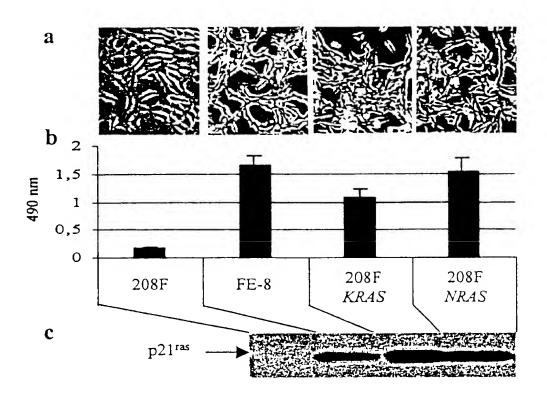


FIG. 9

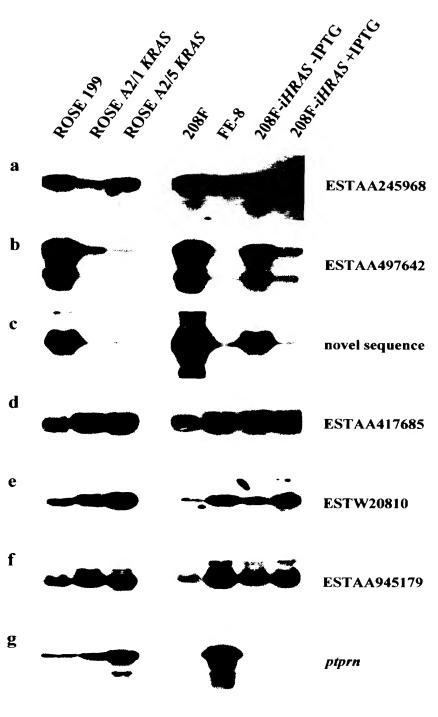


FIG. 10

```
1
2
     T59
     T182
3
     T32
     T6
4
5
     T34
6
7
     N5
     NZC
8
     N280
9
10
11
     N271
      N126
      T148
12
      N199
13
      T64
14
      N131
15
      T20
16
      T162
17
18
      T141
      N77
19
20
      N1C4
      T49
21
      716
22
      N189
23
      N28
24
      T124
      T216
25
26
      T60
27
      T37
28
      T160
29
      N101
30
      N40
31
      T54
32
      T120
33
      N159
34
      T185
35
      N151
36
      T147
37
     N183
38
      T25
39
      T47
40
      T43
41
      T139
42
      T176
43
      N144
44
      T35
45
      T98
      T15
46
     T138
47
48
     N21
49
      T76
     T103
50
51
52
     T44
53
     N31
54
     T243
55
     N129
56
     T193
57
      T132
58
      T137
59
     T217
60
     T191
61
     N42
62
      T156
63
      T67
```

FIG. 11

```
64
     N196
65
     T21
66
     N34
67
     N134
     T119
68
69
     N36
70
     N2C9
7:
     N256
72
     T105
73
     T75
     T153
T169
74
75
76
     T86
77
     T111
73
      T144
79
     N192
80
     N103
     N270
81
     N255
82
83
     NGI
84
     N137
      T174
85
86
     N22
87
      T2
88
      T237
89
      T19
      N156
90
91
     N59
92
      N235
92
     N248
92
     N249
92
     N252
92
      N257
93
      8ET
94
      T121
      Nlo
95
96
      T129
97
      T66
98
      T36
99
      T40
100
      Nl
101
       N212
102
       T100
103
       N112
104
       NЗ
105
       N238
106
       T183
107
       T238
108
       T166
109
       N29
       T225
110
111
       N175
112
       N142
113
       T72
       N186
114
       T212
115
116
       T196
117
       T48
113
       N132
119
       N158
120
       T69
121
       N7
       T245
122
```

**FIG. 11A** 

```
N102
123
      T208
124
      N44
125
125
      T205
      T215
127
      N293
128
123
      T226
130
      T253
131
      T222
132
      N264
133
134
      T240
      N70
135
      T125
136
      N253
137
      N234
138
      N55
139
      NZOZ
140
141
142
      N82
      T45
      T118
143
      710
144
      N71
145
      N183
      N165
146
147
      N213
      N35
148
149
      N182
150
      ,143
151
      N75
       T163
152
153
      T89
154
      N11
155
      N32
156
      T50
157
      N215
158
      N242
159
      N181
160
      N48
161
       T227
162
      N149
      N109
163
164
      N260
165
       T219
166
      T61
167
      N85
168
       N45
169
       T250
170
      N261
171
      T172
172
      N62
173
      N160
174
      N154
175
       B58
176
       T232
177
       N128
178
       N79
179
       T58
130
       N30
131
       T68
      T244
182
192
       T251
132
       T96
```

183

N26

**FIG. 11B** 

```
184
      N14
185
      N121
136
      T17
187
      ΤЗ
133
      T117
139
      T14
      T73
190
191
      N 4
192
      N289
193
      T239
      T170
194
195
      T146
196
      N17
197
      T235
193
      N74
199
      SIN
200
      T211
201
      T136
201
      T204
202
      N50
      N116
203
      T223
204
205
      N198
206
      N267
      T133
207
208
      T80
209
      изів
210
      N266
211
      T224
212
      N143
213
      NIO3
214
      N263
215
      N250
216
      N92
217
      N152
213
      Tll
      T159
219
220
      N243
221
      N78
      T116
222
223
      T27
224
      N207
      ТЗI
225
226
      изв
227
      N163
223
      NSl
229
      T94
230
      N228
231
      И8О
232
      T230
233
      T188
234
      N190
      N187
235
236
      N136
      N294
237
      N275
238
239
      N65
240
      N89
241
      N125
242
      N205
243
      N39
244
      N13
245
       T48
```

FIG. 11C

```
246
       Tloo
247
       T223
248
       N104
249
       N35
250
       T245
251
       N32
252
       T62
253
       N125
254
       N130
255
       N22
256
       T61
       T125
257
       T174
T36
258
259
       T19
260
261
       T204
262
       T153
263
       T27
264
       T212
265
       T159
266
       T226
257
       T239
268
       N263
269
       T66
270
       N75
271
       N250
272
       T175
273
       N283
274
       T40
275
       N152
276
       N256
277
       M28
278
       T160
279
       TE2
       N122
T170
280
281
282
       N44
283
       NIS
284
       T103
285
       N126
236
       N55
287
       T42
288
       T34
289
      N159
290
       N21
291
      N154
292
       N80
       T189
T17
293
294
295
       T68
296
       T14
297
      T146
298
      T120
299
      N181
300
      N192
301
      T109
302
      N215
303
      T244
303
      T251
304
      T96
305
       T211
306
      T243
307
      N213
```

**FIG. 11D** 

```
308
      T224
309
      T94
310
      T183
      N294
T191
311
312
313
      T88
314
      79
315
      N204
316
      N175
317
      N129
318
      T141
319
      N188
320
      N209
321
      T111
322
      T144
323
      N213
324
      N109
325
      N62
326
      T235
327
      N193
328
      N148
329
      N79
330
      T116
331
      N46
332
      N49
333
      N51
334
      N52
335
      T26
```

FIG. 11E